Anticancer Potential of an Ethanol Extract of *Saussurea involucrata* against Hepatic Cancer Cells in *vitro*

Munkhzaya Byambaragchaa¹, Joseph dela Cruz², Altantsetseg Kh³, Seong-Gu Hwang¹*

Abstract

*Saussurea involucrata* is a Mongolian medicinal plant well known for its effects in promoting blood circulation, and anti-inflammation and analgesic functions. Earlier studies reported that *Saussurea involucrata* has anti-cancer activity. The purpose of this study was to confirm the anticancer activity of an ethanol extract of *Saussurea involucrata* against hepatic cancer and elucidate its mechanisms of action. Hepatocellular carcinoma cells were tested in vitro for cytotoxicity, AO/EB staining for apoptotic cells, apoptotic DNA fragmentation and cell cycle distribution in response to *Saussurea involucrata* extract (SIE). The mRNA expression of caspase-3,-9 and Cdk2 and protein expression of caspase-3,-9, PARP, XIAP, Cdk2 and p21 were analyzed through real time PCR and Western blotting. Treatment with SIE inhibited HepG2 cell proliferation dose- and time-dependently, but SIE only exerted a modest cytotoxic effect on viability of Chang human liver cells. Cells exposed to SIE showed typical hallmarks of apoptotic cell death. Cell cycle analysis revealed that SIE caused G1-phase arrest in HepG2 cells. In conclusion, *Saussurea involucrata* ethanol extract has potential cytotoxic and apoptotic effects on human hepatocellular carcinoma cells. Its mechanism of action might be associated with the inhibition of DNA synthesis, cell cycle (G1) arrest and apoptosis induction through up-regulation of the protein expressions of caspase-3,-9 and p21, degradation of PARP and down-regulation of the protein expression of Cdk2 and XIAP.

Keywords: *Saussurea involucrata* - G1 phase arrest - apoptosis - HepG2 cells - growth inhibition

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Introduction

Liver cancer remains the sixth most frequently diagnosed cancer and the third leading cause of cancer-related mortality globally (Chan and Yeo, 2012). It is estimated that 748,300 new liver cancer cases and 695,900 cancer deaths occurred in 2008. The world Health Organization (WHO) estimates that 84 million people would die of cancer between 2005 and 2015. (Jemal et al., 2011). The current trend of cancer research is the investigation of medicines of plant origin because of their affordability and accessibility with minimal side effects. Treatment of hepatocellular carcinoma is clinically difficult, as HCC (hepatocellular carcinoma) expresses multidrug resistance genes and is highly associated with multi-gene, multi-factor, and multi-step processes (Huang et al., 1992; Padhya et al., 2013). HCC cells are characterized by high cellular levels of cyclin dependent kinases (Cdks). Up-regulation of Cdks results from the inactivation of p16⁰ink⁴, p21⁰waf¹ and p27⁰kip¹. Cdk inhibitory proteins, and abnormal activation of cyclins.

*Saussurea involucrata* KAR. et KIR., belongs to the Asteraceae family, is a rare and beneficial traditional medicinal herb, grows in the mountains at heights of 4000-4300m in Mongolia and China. In recent years, the wild sources of *S.involucrata* have decreased dramatically due to the exhaustive collection for use in pharmaceutical preparations. In order to conserve the natural sources of *S.involucrata*, tissue culture is being developed, which might be used as a potential substitute for wild *S.involucrata* in the pharmaceutical industry (Guo et al., 2007; Desai et al., 2008). Cultured and wild *S.involucrata* showed the same phytochemical profile (Guo et al., 2007). The dried aerial parts of *S.involucrata* have long been used for the treatment of rheumatoid arthritis, impotence, irregular menses, cough with cold, stomachache, and altitude sickness (Li et al., 1989; Jia et al., 2005). Modern pharmacological studies showed that this plant has anti-inflammatory and analgesic (Jia et al., 2005), anti-metastasis (Byambaragchaa et al., 2013) and anti cancer (Way et al., 2010) effects. Recent studies have shown that *S.involucrata* extract (SIE) has anti-cancer effect, however information on its potential mechanism remains limited. *S.involucrata* contains anti-cancer compounds such as rutin, hispidulin (Liu et al., 1985), apigenin (Li et al., 2006).
Preparation of SIE
Snow lotus (Saussurea involucrata) collected in Mongolia was used in this study. 10 g air-dried plants were kept in 500 ml 95% ethanol for 48 h and then filtered. The extract was evaporated to dryness with a rotary vacuum evaporator and freeze-dried to give a powder. The stock solution of Saussurea involucrata extract (SIE) was prepared by dissolving snow lotus powder in culture medium (10 mg/mL) and the experimental concentrations were diluted in the basal medium.

Cell culture
Human hepatocarcinoma HepG2 cells were maintained in RPMI-1640 (Gibco) supplemented with 10% heat-inactivated fetal bovine serum (FBS, Gibco), 100 units/mL penicillin, and 100 µg/mL streptomycin under 5% CO₂ in a humidified incubator at 37°C.

Cell viability analysis
Cell viability was quantified by CCK-8 assay. Briefly, the cells were plated in 96-well culture plates at a density of 10⁴ cells/mL and allowed to adhere at 37°C for 12 h. Then cells were exposed to various doses of SIE. At various times after exposure, the media with various SIE concentrations were exchanged; the cells were washed with PBS once. Then fresh media with CCK-8 reagent were added to each well. After 1 hour at 37°C, the absorbance of OD₄₅₀ was determined using an ELISA plate reader (Tecan, Switzerland). Morphological changes after exposure to SIE were observed by an inverted microscope.

Nuclear staining
Cells were exposed to 0-400 µg/mL of SIE for 48 h and stained nuclei with EB/AO dye mix containing 100 µg/mL acridine orange and 100 µg/mL ethidium bromide. Cells were viewed using confocal fluorescent microscope.

Cell cycle analysis
To determine cell cycle distribution analysis, 5×10⁴ cells/mL were plated in 6 well plates and treated with SIE (0-400 µg/mL) for 24 h. After treatment, the cells were collected by trypsinization, fixed in 70% ethanol, washed in PBS, resuspended in 1 mL of PBS containing 1 mg/mL RNase and 50 mg/mL PI, incubated in the dark for 30 min at room temperature, and analyzed by BD FACS Calibur flow cytometer. The data were analyzed using the CellQuestPro software.

DNA fragmentation assay
Cells (5×10⁴/mL) were treated with 0-400 µg/mL of SIE for 24 h, and then cells were collected by centrifugation. Pellets were lysed by DNA lysis buffer (10 mM Tris-HCl, 10 mM EDTA, 0.5% TritonX-100) and then centrifuged. The supernatant obtained was incubated overnight with proteinase K (0.1 mg/mL) and then with RNase (0.2 mg/mL) for 2 hr at 37°C. After extraction with phenol:chloroform:isoamylalcohol (25:24:1), the DNA was separated in 1.8% agarose gel and visualized by UV after staining with ethidium bromide.

Quantitative real time PCR
Total RNA was extracted using Trizol reagent according to the manufacturer’s instructions (Invitrogen, MA, USA). Complementary DNA (cDNA) was obtained by following the protocol of THERMOScript RT-PCR System (Life Technologies). PCR was applied to cDNAs prepared from untreated and SIE treated samples. Quantitative PCR was performed using the Power SYBR Green PCR Master Mix from Applied Biosystems in a Bio-Rad C1000TM Thermal Cycler following the manufacturer’s protocol. Amplification was carried out in a total volume of 20 µL for 40 cycles of 15 s at 95°C, 20 s at 60°C, and 30 s at 72°C. Samples were run in triplicate and their relative expression was determined by normalizing expression of each target GAPDH. These were then compared with the normalized expression in control untreated sample to calculate a change value (%). Primer sequences were as follows: human GAPDH: sense 5'-CGAGATCCCTCCAAATAATCA-3'; antisense 5'-AGGTCCACCCTGACACCGTT-3', human caspase-3: sense 5'-TTTTTCAAGGGGATCTGTG-3', antisense 5'-CGGGTAAAAACGGAATGAT-3', human caspase-9: sense 5'-AGGAACTCAGCAGCTCATTAC-3', antisense 5'-CAGCATTAGCCAGACCTAAGCA-3', human cdk2: sense 5'-GCCCTAAATCTACCACTTCCC-3'; antisense 5'-AAGGTTGTTGAGGGCTAACT-3'.

Western blot
The cells treated with 0-400 µg/mL SIE for 24 h were lysed in a protein extraction solution (Intron Biotechnology). The protein concentration was determined by the Bio-Rad protein assay (Bio-Rad Laboratories, Richmond, CA). For Western blotting, 30 micrograms of protein extraction was separated on 8-15% SDS-
PAGE and electrotransferred to nitrocellulose membrane (Schleicher and Schuell, Germany). The membrane was blocked by incubation in 5% skim milk in TBST buffer (20mM Tris-HCl, pH 7.6, 140mM NaCl, 0.1% Tween 20) and then incubated with monoclonal anti-p21, anti-cdk2, anti-actin (Abcam), monoclonal anti-caspase-3, anti-caspase-9, anti-XIAP, and anti-PARP (Cell signaling technology) for 13-15h. The blot was washed with TBST buffer and incubated with horse radish peroxidase-labeled secondary antibody (Santa Cruz Biotechnology) for 2 h. The membranes were then washed again, and detection was performed using the enhanced chemiluminescence system (Amersham).

Statistical analysis

All in vitro experiments were done in triplicate, and the experiments were repeated at least thrice. The intensities of the mRNA bands were normalized to the GAPDH bands and quantified by comparing with those of control cells. Data were expressed as means ± standard deviations. The difference between control and SIE-treated cells was considered statistically significant.

Results

Inhibition of cell proliferation

First, we tested the effects of SIE on the cell growth of hepatocarcinoma cell line, HepG2 and a normal hepatocyte, Chang normal cell. The cells were exposed to various doses of SIE for 0-72 h and cell viability was determined by CCK-8 assay. As shown in Figure 1 A and B, the survival curve showed that SIE had dose- and time-dependent cytotoxic effects on HepG2 cells. In contrast, Chang normal liver cells were not markedly affected by the treatment with SIE (400 μg/mL) (Figure 1 C, D). SIE reduced the proliferation of HepG2 and Chang liver cells dose- and time- dependently with the IC_{50} value of 84.8μg/mL and 616.99μg/mL, respectively. The inhibition of the cell proliferation could be the result of induction of apoptosis or cell cycle arrest. We hypothesized that SIE blocked hepatocarcinoma cell proliferation which caused the alterations of cell cycle progression and apoptosis.

Changes in nuclei morphology after SIE treatment

To determine whether the growth inhibitory activity of SIE was related to the induction of apoptosis, morphological assay of cell death was investigated using the AO/EB staining for fluorescence microscopy. After HepG2 cells were exposed to various concentrations of SIE for 24 h, different morphological features were analyzed. Uniformly green live cells with normal morphology were seen in the control group (Figure 2A), whereas yellow early apoptotic cells with chromatin condensation and orange later apoptotic cells with fragmented chromatin and apoptotic bodies were seen when SIE was at the concentration of 200 and 400 μg/mL (Figure 2B and C). The results suggested that SIE was able to induce marked apoptotic morphology in HepG2 cells.

Effect of SIE on DNA fragmentation in HepG2 cells

One of the mechanisms by which cell growth is suppressed is apoptotic cell death. Therefore, the effect of SIE on DNA fragmentation was examined in HepG2 cells. The DNA fragmentation was observed when cells were treated with 100, 200, 400 μg/mL of SIE for 24h. The profile for SIE induced apoptosis closely correlated with its growth suppressive effects. Thus, the growth suppression induced by SIE in HepG2 cells may be related to the induction of apoptosis.

Cell cycle regulation

The mechanism responsible for SIE-mediated cell proliferation inhibition, we analyzed the cell cycle

Figure 2. Effect of SIE on the induction of apoptotic morphological changes on HepG2 cells stained with AO/EB. A) Green live cells show normal morphology in control; B) yellow early apoptotic cells show chromatin condensation with 200μg/mL SIE; C) 400μg/mL SIE treatment showed orange later apoptotic cells, fragmented chromatin and apoptotic bodies (×200)

Figure 3. The Effect of SIE Treatment on DNA Fragmentation; Confirmed by Staining with Ethidium Bromide in 1.8% Agarose Gel
progression of HepG2 cells after treatment with SIE (various concentrations) for 24 hours (Figure 4) by flow cytometry. As shown in Figure 4, the results indicated that, compared with the control, 400 µg/mL SIE increased the population of SubG1 (apoptotic cells) phase from 3.74±0.04 to 6.69±0.16% (Figure 4, p<0.001), G1 phase from 68.36±4.03 to 83.66±3.28% (Figure 4, p<0.01) and decreased the population of S phase from 7.34±0.29 to 1.94±0.24 (Figure 4, p<0.001) and G2/M phase from 20.91±4.13 to 7.81±0.6% (Figure 4, p<0.01). Thus, SIE induces cell cycle arrest and apoptosis in HepG2 cells.

**Cell cycle and apoptosis related gene expressions**

SIE induced cell cycle arrest and apoptosis cascade activation was determined further by the mRNA expressions of cell cycle and apoptosis related genes with Real Time PCR. The mRNA expressions of caspase-3 and caspase-9 were up-regulated by 111.5% (p<0.001) and 64.28% (p<0.01), respectively, while the mRNA expression of Cdk2 was down-regulated by 65.32% (p<0.001) after treatment of 400 µg/mL SIE for 12h (Figure 5).

**Cell cycle and apoptosis related protein expressions**

To explore the potential signaling pathways underlying the SIE-induced apoptosis of cancer cells, cell cycle and apoptosis related protein expressions, cleaved caspase-3, cleaved caspase-9, cleaved PARP, XIAP, cdk2 and p21 were evaluated with western blotting. The expression levels of cleaved caspase-3, cleaved caspase-9, cleaved PARP and p21 were increased in dose-dependent manner, and the expression levels of Cdk2 and XIAP were decreased in dose-dependent manner (Figure 6).

**Discussion**

Hepatocellular carcinoma is the most malignant cancer with a high incidence and mortality, for which current therapeutic approaches are still very limited, although chemotherapy is one of the important therapeutic modalities for HCC, the toxic effects are difficult to tolerate (Li et al., 2009; Lu et al., 2010; Roomi et al., 2010; Dai et al., 2011), so, there is an urgent need for development of better drugs which are more effective and simultaneously cause fewer side effects (Mukherjee et al., 2011; Shynu et al., 2011; Li-Weber, 2013). Natural therapies, such as the use of plant-derived products and traditional Chinese medicine (TCM) in cancer treatment, may reduce adverse side effects (Dai et al., 2011; Shynu et al., 2011; Wang et al., 2012), medicinal plants and natural products of plant origin are a rich source of cancer chemotherapy drugs, and exhibited low or almost no toxicity to normal tissues, hence, more attention is being paid to searching for new antitumor agents from natural products (Sigstedt et al., 2008; George et al., 2010).

*Saussurea involucrata* Kar et Kir belongs to the Asteraceae family, is a rare and beneficial traditional medicinal herb in Mongolia and China. The dried aerial parts of *S. involucrata* have long been used for the treatment of common diseases. Recent studies have shown that *S. involucrata* extract (SIE) has anti-cancer effect; however information on its potential mechanism remains limited. Therefore, further studies are needed to clarify the potential anticancer activity of *Saussurea involucrata* and its action mechanism.

In this study, SIE’s anti-proliferative activities on hepatocarcinoma HepG2 cells and human Chang liver normal cells were elucidated. An extract demonstrated significant toxicity on HepG2 with $IC_{50}$ value of at 84.8µg/mL and lower cytotoxicity on Chang liver normal cells with $IC_{50}$ value of 616.99µg/mL.

Cell cycle arrest and apoptosis are two main ways for cell growth inhibition (Sun et al., 2011), natural products primarily target rapidly cycling tumor cells, inducing apoptosis of cancer cells as anticancer agents (Sreelatha et al., 2011).

After treatment with SIE, the characteristics of apoptotic cells, including the increase of EB/AO staining apoptotic cells (Figure 2) and evident DNA fragmentations...
In our result, treatment with SIE caused an induction of -9 can cleave poly-(ADP-ribose) polymerase (PARP). Proteolysis during apoptosis. For example, caspase-3 and cleavage of cellular proteins which are characteristically by removal of the prodomain and cleavage between procaspases, with the active tetramer being formed proteases). The caspases are present in cells as inactive execution of apoptotic signals requires the coordinated cytochrome-c pathway by inhibiting caspase-9 (Deveraux al., 2006). Inhibitor of apoptosis protein (IAP) is a family ability to block cells from undergoing apoptosis (Dong et al., 2006). Overexpression of the tumor suppressor gene products, is known to be one of the critical events that drive cancer cells into uncontrolled proliferation (Dong et al., 2006). Cdk2 orchestrates the orderly progression of the eukaryotic cell cycle and plays a key role in the progression from late G1 to late G2 phase (Van den Heuvel and Harlow, 1993). It has been suggested that prolonged inhibition of Cdk2 within cells induces apoptosis and studies have speculated that Cdk2 inhibition decreases XIAP expression, releasing their ability to block cells from undergoing apoptosis (Dong et al., 2006). Inhibitor of apoptosis protein (IAP) is a family of proteins that regulate cell death; X-linked IAP (XIAP) is important member of this family in mammals. It inhibits caspasas (caspase-3 and caspase-9) and block apoptosis (Salvesen and Duckett, 2002). The apoptotic effects of SIE in HepG2 cells appear to relate to changes in the cellular level of anti-apoptotic protein. Figure 6 shows that protein level of Cdk2 and XIAP are down-regulated by SIE in a dose-dependent manner in HepG2 cells. This finding suggests that increased p21 is involved in cell cycle arrest by SIE.

Cdk2 regulate two biological processes essential for cancer cell survival; cell cycle progression and gene transcription (Nurse, 2009). Deregulation of the cell cycle, a process controlled by various cyclins, cyclin-dependent kinases (CDKs), CDK inhibitors, and certain tumor suppressor gene products, is known to be one of the critical events that drive cancer cells into uncontrolled proliferation. Generally, cell cycle arrest and apoptosis are connected; occurrence of cell cycle arrest leads to cell apoptosis, which need numerous signaling molecules and regulatory proteins. p21 plays an essential role in growth arrest after DNA damage, overexpression leads to G1 and G2 arrest and is highlighted as an important tumor suppressor (Erol, 2011; Masgras et al., 2012). It can bind and inhibit a broad range of cyclin/Cdk complexes, with preference for those containing Cdk2 (Zhang et al., 1993).

Figure 6 shows that protein level of p21 is up-regulated by SIE in a dose-dependent manner in HepG2 cells. This finding suggests that increased p21 is involved in cell cycle arrest by SIE.

Taken together, the potential anti-cancer activity of the SIE against hepatic cancer in vitro and its partial molecular mechanisms of activities were investigated in this experimental study. The results demonstrated that the Saussurea involucrata extract has strong anti-cancer activity against liver cancer without significant effect on normal cells. And the mechanism of its action might be associated with the inhibition of DNA synthesis, cell cycle arrest in G1 phase, apoptosis induction through caspase-3-mediated pathway. The Saussurea involucrata is a strong candidate for use an anti-cancer therapeutic agent for the treatment of hepatocellular carcinoma, and deserve to be investigated further.

References


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Molecules

Current Drug Metabolism

EMBO J

Cancer Biol Ther

Cancer Prev Res (癌研究)

J Gastroenterol Hepatol

Cell Cycle

MCC Complement Altern Med

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